

SEQUENCE LISTING

<110> KANNOUCHE, PATRICIA

MAUFFREY, PHILLIPE

PINON-LATAILLADE, GHISLAINE

BIARD, DENIS

ANGULO-MORA, JAINE

<120> SEQUENCES CODING FOR KIN17 PROTEIN AND THEIR APPLICATIONS

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<140> 09/555,529

<141> 2000-07-24

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<151> 1998-12-09

<150> FR97 15536

<151> 1997-12-09

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<170> PatentIn version 3.0

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caacaacatt gtctacaacg aatacatcag ccaccgagag cacatccaca tgaatgccac	180
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Lys Gln Cys Arg Asp Glu Asn Gly Phe Lys Cys His Cys Met Ser Glu  
35 40 45

Ser His Gln Arg Gln Leu Leu Leu Ala Ser Glu Asn Pro Gln Gln Phe  
50 55 60

Met Asp Tyr Phe Ser Glu Glu Phe Arg Asn Asp Phe Leu Glu Leu Leu  
65 70 75 80

Arg Arg Arg Phe Gly Thr Lys Arg Val His Asn Asn Ile Val Tyr Asn  
85 90 95

Glu Tyr Ile Ser His Arg Glu His Ile His Met Asn Ala Thr Gln Trp  
100 105 110

Glu Thr Leu Thr Asp Phe Thr Lys Trp Leu Gly Arg Glu Gly Leu Cys  
115 120 125

Ala Leu Lys Leu Leu Gly Ser Ala Ala Ser Gly Lys Arg Lys Glu Ser  
130 135 140

Ser Gln Ser Ser Ala Gln Pro Ala Lys Lys Lys Ser Ala Leu Asp  
145 150 155 160

Glu Ile Met Glu Leu Glu Glu Lys Lys Arg Thr Ala Arg Thr Asp  
165 170 175

Ala Trp Leu Gln Pro Gly Ile Val Val Lys Ile Ile Thr Lys Lys Leu  
180 185 190

Gly Glu Lys Tyr His Lys Lys Gly Val Val Lys Glu Val Ile Asp  
195 200 205

Arg Tyr Thr Ala Val Val Lys Met Thr Asp Ser Gly Asp Arg Leu Lys  
210 215 220

Leu Asp Gln Thr His Leu Glu Thr Val Ile Pro Ala Pro Gly Lys Arg  
225 230 235 240

Val Leu Val Leu Asn Gly Gly Tyr Arg Gly Asn Glu Gly Thr Leu Glu  
245 250 255

Ser Ile Asn Glu Lys Ala Phe Ser Ala Thr Ile Val Ile Glu Thr Gly  
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Lys Leu Ala  
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<213> Homo sapiens

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Lys Gln Cys Arg Asp Glu Asn Gly Phe Lys Cys His Cys Met Ser Glu  
35 40 45

Ser His Gln Arg Gln Leu Leu Ala Ser Glu Asn Pro Gln Gln Phe  
50 55 60

Met Asp Tyr Phe Ser Glu Glu Phe Arg Asn Asp Phe Leu Glu Leu Leu  
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Arg Arg Arg Phe Gly Thr Lys Arg Val His Asn Asn Ile Val Tyr Asn  
85 90 95

Glu Tyr Ile Ser His Arg Glu His Ile His Met Asn Ala Thr Gln Trp  
100 105 110

Glu Thr Leu Thr Asp Phe Thr Lys Trp Leu Gly Arg Glu Gly Leu Cys  
115 120 125

Ala Leu Lys Thr Ile Gly Ser Ser Ala Ser Val Lys Arg Lys Glu Ser  
130 135 140

Ser Gln Ser Ser Thr Gln Ser Lys Glu Lys Lys Lys Lys Ser Ala  
145 150 155 160

Leu Asp Glu Ile Met Glu Ile Glu Glu Lys Lys Arg Thr Ala Arg  
165 170 175

Thr Asp Tyr Trp Leu Gln Pro Glu Ile Ile Val Lys Ile Ile Thr Lys  
180 185 190

Lys Leu Gly Glu Lys Tyr His Lys Lys Lys Ala Ile Val Lys Glu Val  
195 200 205

Ile Asp Lys Tyr Thr Ala Val Val Lys Met Ile Asp Ser Gly Asp Lys  
210 215 220

Leu Lys Leu Asp Gln Thr His Leu Glu Thr Val Ile Pro Ala Pro Gly  
225 230 235 240

Lys Arg Ile Leu Val Leu Asn Gly Gly Tyr Arg Gly Asn Glu Gly Thr  
245 250 255

Leu Glu Ser Ile Asn Glu Lys Thr Phe Ser Ala Thr Ile Val Ile Glu  
260 265 270

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Ile Ser Lys Leu Ala  
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Lys Gln Cys Arg Asp Glu Asn Gly Phe Lys Cys His Cys Met Ser Glu  
35 40 45

Ser His Gln Arg Gln Leu Leu Leu Ala Ser Glu Asn Pro Gln Gln Phe

50 55 60

Met Asp Tyr Phe Ser Glu Glu Phe Arg Asn Asp Phe Leu Glu Leu Leu  
65 70 75 80

Arg Arg Arg Phe Gly Thr Lys Arg Val His Asn Asn Ile Val Tyr Asn  
85 90 95

Glu Tyr Ile Ser His Arg Glu His Ile His Met Asn Ala Thr Gln Trp  
100 105 110

Glu Thr Leu Thr Asp Phe Thr Lys Trp Leu Gly Arg Glu Gly Leu Cys  
115 120 125

Lys Val Asp Glu Thr Pro Lys Gly Trp Tyr Ile Gln Tyr Ile Asp Arg  
130 135 140

Asp Pro Glu Thr Ile Arg Arg Gln Leu Glu Leu Glu Lys Lys Lys Lys  
145 150 155 160

Gln Asp Leu Asp Asp Glu Glu Lys Thr Ala Lys Phe Ile Glu Glu Gln  
165 170 175

Val Arg Arg Gly Leu Glu Gly Lys Glu Gln Glu Thr Pro Val Phe Thr  
180 185 190

Glu Leu Ser Arg Glu Asn Glu Glu Lys Val Thr Phe Asn Leu Asn  
195 200 205

Lys Gly Ala Gly Gly Ser Ala Gly Ala Thr Thr Ser Lys Ser Ser Ser  
210 215 220

Leu Gly Pro Ser Ala Leu Lys Leu Leu Gly Ser Ala Ala Ser Gly Lys  
225 230 235 240

Arg Lys Glu Ser Ser Gln Ser Ser Ala Gln Pro Ala Lys Lys Lys Lys  
245 250 255

Ser Ala Leu Asp Glu Ile Met Glu Leu Glu Glu Lys Lys Arg Thr  
260 265 270

Ala Arg Thr Asp Ala Trp Leu Gln Pro Gly Ile Val Val Lys Ile Ile  
275 280 285

Thr Lys Lys Leu Gly Glu Lys Tyr His Lys Lys Lys Gly Val Val Lys  
290 295 300

Glu Val Ile Asp Arg Tyr Thr Ala Val Val Lys Met Thr Asp Ser Gly  
305 310 315 320

Asp Arg Leu Lys Leu Asp Gln Thr His Leu Glu Thr Val Ile Pro Ala  
325 330 335

Pro Gly Lys Arg Val Leu Val Leu Asn Gly Gly Tyr Arg Gly Asn Glu  
340 345 350

Gly Thr Leu Glu Ser Ile Asn Glu Lys Ala Phe Ser Ala Thr Ile Val  
355 360 365

Ile Glu Thr Gly Pro Leu Lys Gly Arg Arg Val Glu Gly Ile Gln Tyr  
370 375 380

Glu Asp Ile Ser Lys Leu Ala  
385 390

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<211> 393

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<213> Homo sapiens

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Lys Gln Cys Arg Asp Glu Asn Gly Phe Lys Cys His Cys Met Ser Glu  
35 40 45

Ser His Gln Arg Gln Leu Leu Ala Ser Glu Asn Pro Gln Gln Phe  
50 55 60

Met Asp Tyr Phe Ser Glu Glu Phe Arg Asn Asp Phe Leu Glu Leu Leu  
65 70 75 80

Arg Arg Arg Phe Gly Thr Lys Arg Val His Asn Asn Ile Val Tyr Asn  
85 90 95

Glu Tyr Ile Ser His Arg Glu His Ile His Met Asn Ala Thr Gln Trp  
100 105 110

Glu Thr Leu Thr Asp Phe Thr Lys Trp Leu Gly Arg Glu Gly Leu Cys  
115 120 125

Lys Val Asp Glu Thr Pro Lys Gly Trp Tyr Ile Gln Tyr Ile Asp Arg  
130 135 140

Asp Pro Glu Thr Ile Arg Arg Gln Leu Glu Leu Glu Lys Lys Lys Lys  
145 150 155 160

Gln Asp Leu Asp Asp Glu Glu Lys Thr Ala Lys Phe Ile Glu Glu Gln  
165 170 175

Val Arg Arg Gly Leu Glu Gly Lys Glu Gln Glu Val Pro Thr Phe Thr  
180 185 190

Glu Leu Ser Arg Glu Asn Asp Glu Glu Lys Val Thr Phe Asn Leu Ser  
195 200 205

Lys Gly Ala Cys Ser Ser Ser Gly Ala Thr Ser Ser Lys Ser Ser Thr  
210 215 220

Leu Gly Pro Ser Ala Leu Lys Thr Ile Gly Ser Ser Ala Ser Val Lys  
225 230 235 240

Arg Lys Glu Ser Ser Gln Ser Ser Thr Gln Ser Lys Glu Lys Lys Lys  
245 250 255

Lys Lys Ser Ala Leu Asp Glu Ile Met Glu Ile Glu Glu Glu Lys Lys  
260 265 270

Arg Thr Ala Arg Thr Asp Tyr Trp Leu Gln Pro Glu Ile Ile Val Lys  
275 280 285

Ile Ile Thr Lys Lys Leu Gly Glu Lys Tyr His Lys Lys Lys Ala Ile  
290 295 300

Val Lys Glu Val Ile Asp Lys Tyr Thr Ala Val Val Lys Met Ile Asp  
305 310 315 320

Ser Gly Asp Lys Leu Lys Leu Asp Gln Thr His Leu Glu Thr Val Ile  
325 330 335

Pro Ala Pro Gly Lys Arg Ile Leu Val Leu Asn Gly Gly Tyr Arg Gly  
340 345 350

Asn Glu Gly Thr Leu Glu Ser Ile Asn Glu Lys Thr Phe Ser Ala Thr  
355 360 365

Ile Val Ile Glu Thr Gly Pro Leu Lys Gly Arg Arg Val Glu Gly Ile  
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Gln Tyr Glu Asp Ile Ser Lys Leu Ala  
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<400> 27

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<212> PRT

<213> Homo sapiens

<400> 35

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Glu Gly Leu Tyr Gly Leu Leu Tyr Ser Thr Tyr Arg His Ile Ser Leu  
35 40 45

Tyr Ser Leu Tyr Ser Leu Tyr Ser Ala Leu Ala Ile Leu Glu Val Ala  
50 55 60

Leu Leu Tyr Ser Gly Leu Val Ala Leu Ile Leu Glu Ala Ser Pro Leu  
65 70 75 80

Tyr Ser Thr Tyr Arg Thr His Arg Ala Leu Ala Val Ala Leu Val Ala  
85 90 95

Leu Leu Tyr Ser Met Glu Thr Ile Leu Glu Ala Ser Pro Ser Glu Arg  
100 105 110

Gly Leu Tyr Ala Ser Pro Leu Tyr Ser Leu Glu Leu Tyr Ser Leu Glu  
115 120 125

Ala Ser Pro Gly Leu Asn Thr His Arg His Ile Ser Leu Glu Gly Leu  
130 135 140

Thr His Arg Val Ala Leu Ile Leu Glu Pro Arg Ala Leu Ala Pro Arg  
145 150 155 160

Gly Leu Tyr Leu Tyr Ser Ala Arg Gly Ile Leu Glu Leu Glu Val Ala  
165 170 175

Leu Leu Glu Ala Ser Asn Gly Leu Tyr Gly Leu Tyr Thr Tyr Arg Ala  
180 185 190

Arg Gly Gly Leu Tyr Ala Ser Asn Gly Leu Gly Leu Tyr Thr His Arg  
195 200 205

Leu Glu Gly Leu Ser Glu Arg Ile Leu Glu Ala Ser Asn Gly Leu Leu  
210 215 220

Tyr Ser Thr His Arg Pro His Glu Ser Glu Arg Ala Leu Ala Thr His  
225 230 235 240

Arg Ile Leu Glu Val Ala Leu Ile Leu Glu Gly Leu Thr His Arg Gly  
245 250 255

Leu Tyr Pro Arg Leu Glu Leu Tyr Ser Gly Leu Tyr Ala Arg Gly Ala  
260 265 270

Arg Gly Val Ala Leu Gly Leu Gly Leu Tyr Ile Leu Glu Gly Leu Asn  
275 280 285

Thr Tyr Arg Gly Leu Ala Ser Pro Ile Leu Glu Ser Glu Arg Leu Tyr  
290 295 300

Ser Leu Glu Ala Leu Ala  
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<210> 36

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<212> PRT

<213> Mus sp.

<400> 36

Pro Arg Gly Leu Tyr Ile Leu Glu Val Ala Leu Val Ala Leu Leu Tyr  
1 5 10 15

Ser Ile Leu Glu Ile Leu Glu Thr His Arg Leu Tyr Ser Leu Tyr Ser  
20 25 30

Leu Glu Gly Leu Tyr Gly Leu Leu Tyr Ser Thr Tyr Arg His Ile Ser  
35 40 45

Leu Tyr Ser Leu Tyr Ser Leu Tyr Ser Gly Leu Tyr Val Ala Leu Val  
50 55 60

Ala Leu Leu Tyr Ser Gly Leu Val Ala Leu Ile Leu Glu Ala Ser Pro  
65 70 75 80

Ala Arg Gly Thr Tyr Arg Thr His Arg Ala Leu Ala Val Ala Leu Val  
85 90 95

Ala Leu Leu Tyr Ser Met Glu Thr Thr His Arg Ala Ser Pro Ser Glu  
100 105 110

Arg Gly Leu Tyr Ala Ser Pro Ala Arg Gly Leu Glu Leu Tyr Ser Leu  
115 120 125

Glu Ala Ser Pro Gly Leu Asn Thr His Arg His Ile Ser Leu Glu Gly  
130 135 140

Leu Thr His Arg Val Ala Leu Ile Leu Glu Pro Arg Ala Leu Ala Pro  
145 150 155 160

Arg Gly Leu Tyr Leu Tyr Ser Ala Arg Gly Val Ala Leu Leu Glu Val  
165 170 175

Ala Leu Leu Glu Ala Ser Asn Gly Leu Tyr Gly Leu Tyr Thr Tyr Arg  
180 185 190

Ala Arg Gly Gly Leu Tyr Ala Ser Asn Gly Leu Gly Leu Tyr Thr His  
195 200 205

Arg Leu Glu Gly Leu Ser Glu Arg Ile Leu Glu Ala Ser Asn Gly Leu  
210 215 220

Leu Tyr Ser Ala Leu Ala Pro His Glu Ser Glu Arg Ala Leu Ala Thr  
225 230 235 240

His Arg Ile Leu Glu Val Ala Leu Ile Leu Glu Gly Leu Thr His Arg

245

250

255

Gly Leu Tyr Pro Arg Leu Glu Leu Tyr Ser Gly Leu Tyr Ala Arg Gly  
260 265 270

Ala Arg Gly Val Ala Leu Gly Leu Gly Leu Tyr Ile Leu Glu Gly Leu  
275 280 285

*α*  
Asn Thr Tyr Arg Gly Leu Ala Ser Pro Ile Leu Glu Ser Glu Arg Leu  
290 295 300

Tyr Ser Leu Glu Ala Leu Ala  
305 310